

Investigation on interaction between Ligupurpuroside docking methods

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Citation Report

#	ARTICLE	IF	CITATIONS
1	The soluble recombinant N-terminal domain of HMW 1Dx5 and its aggregation behavior. <i>Food Research International</i> , 2015, 78, 201-208.	2.9	23
2	Molecular interactions of flavonoids to pepsin: Insights from spectroscopic and molecular docking studies. <i>Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy</i> , 2015, 151, 576-590.	2.0	39
3	Investigation of the Binding Between Pepsin and Nucleoside Analogs by Spectroscopy and Molecular Simulation. <i>Journal of Fluorescence</i> , 2015, 25, 451-463.	1.3	18
4	Interaction Behavior Between Niclosamide and Pepsin Determined by Spectroscopic and Docking Methods. <i>Journal of Fluorescence</i> , 2015, 25, 1681-1693.	1.3	18
5	Studies on the binding of pepsin with three pyrethroid insecticides by multi-spectroscopic approaches and molecular docking. <i>Journal of Molecular Recognition</i> , 2016, 29, 476-484.	1.1	20
6	Probing the Interaction between Acotiamide Hydrochloride and Pepsin by Multispectral Methods, Electrochemical Measurements, and Docking Studies. <i>Journal of Biochemical and Molecular Toxicology</i> , 2016, 30, 350-359.	1.4	3
7	Study on the interaction of β -carotene and astaxanthin with trypsin and pepsin by spectroscopic techniques. <i>Luminescence</i> , 2016, 31, 782-792.	1.5	23
8	Probing the binding mechanisms of α -tocopherol to trypsin and pepsin using isothermal titration calorimetry, spectroscopic, and molecular modeling methods. <i>Journal of Biological Physics</i> , 2016, 42, 415-434.	0.7	19
9	Influence of CuO nanoparticles and nanographene platelets on the photosonocatalytic performance of Fe ₃ O ₄ /TiO ₂ nanocomposites. <i>Journal of Physics: Conference Series</i> , 2016, 776, 012022.	0.3	3
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12	Probing the binding of procyanidin B3 to trypsin and pepsin: A multi-technique approach. <i>International Journal of Biological Macromolecules</i> , 2016, 85, 168-178.	3.6	14
13	Interaction mechanism of pepsin with a natural inhibitor gastrodin studied by spectroscopic methods and molecular docking. <i>Medicinal Chemistry Research</i> , 2017, 26, 405-413.	1.1	18
14	Probing deep into the binding mechanisms of folic acid with α -amylase, pepsin and trypsin: An experimental and computational study. <i>Food Chemistry</i> , 2017, 226, 128-134.	4.2	32
15	Mechanism and Nature of Inhibition of Trypsin by Ligupurpuroside A, a Ku-Ding Tea Extract, Studied by Spectroscopic and Docking Methods. <i>Food Biophysics</i> , 2017, 12, 78-87.	1.4	22
16	Interaction between azo dye Acid Red 14 and pepsin by multispectral methods and docking studies. <i>Luminescence</i> , 2017, 32, 1123-1130.	1.5	17
17	Characterization and analysis of binding of Thioflavin T with partially folded and native states of α -lactalbumin protein by calorimetric and spectroscopic techniques. <i>International Journal of Biological Macromolecules</i> , 2017, 95, 376-384.	3.6	5
18	Investigation and comparison of the binding between tolvaptan and pepsin and trypsin: Multi-spectroscopic approaches and molecular docking. <i>Journal of Molecular Recognition</i> , 2017, 30, e2598.	1.1	17

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19	Dissecting the Disulfide Linkage of the N-Terminal Domain of HMW 1Dx5 and Its Contributions to Dough Functionality. <i>Journal of Agricultural and Food Chemistry</i> , 2017, 65, 6264-6273.	2.4	21
20	Conformation change of trypsin induced by acteoside as studied using multiple spectroscopic and molecular docking methods. <i>International Journal of Food Properties</i> , 2018, 21, 301-312.	1.3	17
21	Effects of acetazolamide on the conformations and activities of digestive enzymes: pepsin and trypsin. <i>Medicinal Chemistry Research</i> , 2018, 27, 1549-1557.	1.1	2
22	Exploring the binding pattern between pepsin and deferasirox using detailed experimental and computer simulation methods. <i>RSC Advances</i> , 2018, 8, 37208-37218.	1.7	8
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24	Binding mechanism of lipase to Ligupurpuroside B extracted from Ku-Ding tea as studied by multi-spectroscopic and molecular docking methods. <i>International Journal of Biological Macromolecules</i> , 2018, 120, 1345-1352.	3.6	26
25	Interaction between trelagliptin and pepsin through spectroscopy methods and molecular dynamics simulation. <i>Spectroscopy Letters</i> , 2018, 51, 332-339.	0.5	4
26	Exploring inhibition mechanism and nature of lipase by Ligupurpuroside A extracted from Ku-Ding tea. <i>Medicinal Chemistry Research</i> , 2018, 27, 1822-1833.	1.1	10
27	Investigation on the Interaction Behavior Between Oenothain B and Pepsin by Isothermal Titration Calorimetry and Spectral Studies. <i>Journal of Food Science</i> , 2019, 84, 2412-2420.	1.5	4
28	Probing the interaction of pepsin with imidacloprid via DFT calculation, spectroscopic approaches and molecular docking. <i>Journal of Molecular Structure</i> , 2019, 1197, 210-216.	1.8	36
29	Trypsin inhibition by Ligupurpuroside B as studied using spectroscopic, CD, and molecular docking techniques. <i>Journal of Biomolecular Structure and Dynamics</i> , 2019, 37, 3379-3387.	2.0	5
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32	Binding of triclosan and triclocarban to pepsin: DFT, spectroscopic and dynamic simulation studies. <i>Chemosphere</i> , 2019, 214, 278-287.	4.2	27
33	Interactions of indole alkaloids with myoglobin: A mass spectrometry based spectrometric and computational method. <i>Rapid Communications in Mass Spectrometry</i> , 2020, 34, e8656.	0.7	3
34	Spectroscopic and molecular docking studies of the interaction between meloxicam and pepsin. <i>Spectroscopy Letters</i> , 2020, 53, 32-43.	0.5	4
35	Multi spectroscopy and molecular modeling aspects related to drug interaction of aspirin and warfarin with pepsin; structural change and protease activity. <i>Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy</i> , 2020, 228, 117813.	2.0	13
36	Interaction of food-grade titanium dioxide nanoparticles with pepsin in simulated gastric fluid. <i>LWT - Food Science and Technology</i> , 2020, 134, 110208.	2.5	16

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37	Investigations of conformational structures and activities of trypsin and pepsin affected by food colourant allura red. <i>Journal of Molecular Liquids</i> , 2020, 319, 114359.	2.3	14
38	â€œRigidâ€•structure is a key determinant for the low digestibility of myoglobin. <i>Food Chemistry: X</i> , 2020, 7, 100094.	1.8	13
39	Interaction behavior between five flavonoids and pepsin: Spectroscopic analysis and molecular docking. <i>Journal of Molecular Structure</i> , 2021, 1223, 128978.	1.8	18
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42	Interactions and effects of food additive dye Allura red on pepsin structure and protease activity; experimental and computational supports. <i>Research in Pharmaceutical Sciences</i> , 2021, 16, 58.	0.6	2
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44	Investigation on the Molecular and Physicochemical Changes of Protein and Starch of Wheat Flour during Heating. <i>Foods</i> , 2021, 10, 1419.	1.9	1
45	Insights on the interaction mechanism of exemestane to three digestive enzymes by multi-spectroscopy and molecular docking. <i>International Journal of Biological Macromolecules</i> , 2021, 187, 54-65.	3.6	6
46	Investigation on the interaction behavior between safranal and pepsin by spectral and MD simulation studies. <i>Journal of Molecular Liquids</i> , 2021, 344, 117903.	2.3	14
47	Application of Molecular Docking in Studies on the Binding Mechanism of Three Enzymes with Natural Products. <i>Advances in Medical Technologies and Clinical Practice Book Series</i> , 2016, , 81-126.	0.3	0
48	Investigation on detoxication effects of 2-hydroxypropyl-Î²-cyclodextrin over two halogenated aromatic DBPs 2,4,6-trichlorophenol and 2,4,6-tribromophenol binding with human serum albumin. <i>Food Chemistry</i> , 2022, 382, 132349.	4.2	3
49	Inhibitory interaction of narcissoside on Î±-glucosidase from <i>Aspergillus niger</i> and <i>Saccharomyces cerevisiae</i> by spectral analysis and molecular docking. <i>Journal of Molecular Structure</i> , 2022, 1264, 133262.	1.8	7
50	The interactions between Reactive Black 5 and human serum albumin: combined spectroscopic and molecular dynamics simulation approaches. <i>Environmental Science and Pollution Research</i> , 2022, 29, 70114-70124.	2.7	8
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